

Supplementary Table 1. Microsatellite loci used to genotype Finnish *Daphnia magna*

populations. Primer sequences and reaction conditions are given in Colson *et al.* (2009). Loci listed in the same group were amplified and genotyped together. Additionally, groups five and six were mixed with groups three and two (respectively) before genotyping.

Locus	Repeat Unit	Tag	Group
WFes0001526	3 (TTA)	FAM	1
WFes0008397	2 (CA)	FAM	1
WFes0007834	2 (CA)	FAM	1
WFes0008608	3 (CAA)	HEX	1
WFes0002528	3 (GGT)	FAM	2
WFes0008344	2 (CT)	HEX	2
WFes0004775	3 (CAA)	HEX	2
WFes0003617	3 (GTT)	FAM	3
WFes0004208	3 (TGA)	FAM	3
WFes0008711	2 (GT)	FAM	4
WFes0009477	3 (GCT)	HEX	4
WFes0004614	2 (GT)	HEX	4
WFes0009449	3 (GTT)	FAM	5
WFes0007001	2 (CA)	HEX	6

Robinson_SupMat_Table2.doc
Genetic estimates of population age - JHered

Supplementary Table 2: Summary of age estimates in simulated datasets. Means, relative biases, and coefficients of variation were calculated from 200 replicate datasets simulated under each parameter combination

A. Estimated ages for simulated datasets using the P _k Distribution (τ)															
5 loci	Ne = 1000					Ne = 10,000					Ne = 1,000,000				
TRUE	10	25	100	250	1000	10	25	100	250	1000	10	25	100	250	1000
Mean	9.08	31.59	98.82	201.71	726.06	12.89	25.36	96.91	239.68	833.81	12.48	25.94	101.26	222.95	834.42
Relative Bias (%)	-10.13	20.86	-1.19	-23.94	-37.73	22.42	1.42	-3.19	-4.31	-19.93	19.87	3.62	1.24	-12.13	-19.84
CV (%)	237.95	164.12	84.76	76.03	54.41	303.54	85.81	66.58	35.03	26.97	185.13	133.31	42.54	28.31	20.14
10 loci	Ne = 1000					Ne = 10,000					Ne = 1,000,000				
TRUE	10	25	100	250	1000	10	25	100	250	1000	10	25	100	250	1000
Mean	10.03	26.25	102.67	234.94	724.49	9.61	25.95	103.16	234.75	840.91	12.19	28.95	101.01	241.85	839.58
Relative Bias (%)	0.30	4.76	2.60	-6.41	-38.03	-4.06	3.66	3.06	-6.50	-18.92	17.97	13.64	1.00	-3.37	-19.11
CV (%)	107.43	104.73	62.40	50.30	39.75	111.24	76.55	41.79	26.04	17.18	144.6	93.66	39.39	21.57	13.09
20 loci	Ne = 1000					Ne = 10,000					Ne = 1,000,000				
TRUE	10	25	100	250	1000	10	25	100	250	1000	10	25	100	250	1000
Mean	11.05	28.29	99.16	224.61	709.43	13.09	24.88	98.5	239.57	820.51	12.69	28.03	101.45	239.78	832.81
Relative Bias (%)	9.50	11.63	-0.85	-11.30	-40.96	23.61	-0.48	-1.52	-4.35	-21.88	21.20	10.81	1.43	-4.26	-20.08
CV (%)	83.24	72.52	38.74	35.98	28.25	140.48	66.72	28.10	18.27	13.28	132.73	59.64	26.21	15.40	9.85
B. Estimated ages for simulated datasets using the Variance in Allele Size (σ_{AS}^2)															
5 loci	Ne = 1000					Ne = 10,000					Ne = 1,000,000				
TRUE	10	25	100	250	1000	10	25	100	250	1000	10	25	100	250	1000
Mean	9.08	31.62	101.06	211.91	797.28	12.89	25.39	97.37	251.97	965.47	12.5	25.97	102.57	230.24	1011.16
Relative Bias (%)	-10.13	20.94	1.05	-17.97	-25.43	22.42	1.54	-2.70	0.78	-3.58	20.00	3.74	2.51	-8.58	1.10
CV (%)	237.93	164.06	86.28	77.03	60.16	303.51	85.93	67.01	35.95	26.19	185.13	133.32	41.79	29.73	16.82
10 loci	Ne = 1000					Ne = 10,000					Ne = 1,000,000				
TRUE	10	25	100	250	1000	10	25	100	250	1000	10	25	100	250	1000
Mean	10.03	26.77	104.11	244.84	810.51	9.61	26.24	104.6	247.14	979.37	12.2	28.99	102.94	254.34	1006.76
Relative Bias (%)	0.30	6.61	3.95	-2.11	-23.38	-4.06	4.73	4.40	-1.16	-2.11	18.03	13.76	2.86	1.71	0.67
CV (%)	107.49	104.87	63.56	49.73	42.27	111.24	76.78	41.69	26.75	16.92	144.57	93.61	40.48	20.99	12.09
20 loci	Ne = 1000					Ne = 10,000					Ne = 1,000,000				
TRUE	10	25	100	250	1000	10	25	100	250	1000	10	25	100	250	1000
Mean	11.05	28.32	100.37	232.84	804.54	13.09	25.01	100.31	251.81	968.28	12.71	28.07	104.24	253.72	1005.42
Relative Bias (%)	9.50	11.72	0.37	-7.37	-24.29	23.61	0.04	0.31	0.72	-3.28	21.32	10.94	4.07	1.47	0.54
CV (%)	83.24	72.52	39.49	37.54	31.87	140.46	66.69	28.20	18.78	12.93	133.28	59.69	25.85	14.35	7.66